COMPLETE LISTING OF CLAIMS

- 1 (Original) A monoclonal or polyclonal antibody having high affinity for a peptide selected from the group consisting of: Phe-Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:1), Phe-Xaa-Gly-Leu-Met-NH₂, where Xaa is variant Phe or Val (SEQ ID NO:4), and Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:2).
- 2 (Original) The antibody of claim 1, wherein said antibody is a monoclonal antibody.
- 3 (Original) The antibody of claim 1, wherein said antibody has high affinity for the peptide Phe-Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:1).
- 4 (Original) The antibody of claim 2, wherein said antibody has high affinity for the peptide Phe-Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:1).
- 5 (Original) The antibody of claim 1, wherein said antibody has high affinity for the peptide Phe-Xaa-Gly-Leu-Met-NH₂, where Xaa is variant Phe or Val (SEQ ID NO:4).
- 6 (Original) The antibody of claim 2, wherein said antibody has high affinity for the peptide Phe-Xaa-Gly-Leu-Met-NH₂, where Xaa is variant Phe or Val (SEQ ID NO:4).
- 7 (Original) The antibody of claim 1, wherein said antibody has high affinity for the peptide Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:2)).
- 8 (Original) The antibody of claim 2, wherein said antibody has high affinity for the peptide Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:2).
- 9 (Original) The antibody of claim 1, wherein said antibody is cross-reactive with each of peptides Phe-Phe-Gly-Leu-Met-NH $_2$ (SEQ ID NO:1), Phe-Xaa-Gly-Leu-Met-NH $_2$, where Xaa is variant Phe or Val (SEQ ID NO4), and Phe-Gly-Leu-Met-NH $_2$ (SEQ ID NO:2).

- (Original) The antibody of claim 2, wherein said antibody is cross-reactive with each of peptides Phe-Phe-Gly-Leu-Met-NH $_2$ (SEQ ID NO:1), Phe-Xaa-Gly-Leu-Met-NH $_2$, where Xaa is variant Phe or Val (SEQ ID NO:4), and Phe-Gly-Leu-Met-NH $_2$ (SEQ ID NO:2).
- (Currently amended) A method for detecting magnesium binding defect comprising:
 - a) measuring in <u>body fluids other than blood plasma</u> blood serum the level of peptide having an amino acid sequence selected from the group consisting of: Phe-Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:1), Phe-Xaa-Gly-Leu-Met-NH₂, where Xaa is variant Phe or Val (SEQ ID NO:4), and Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:2); and
- b) comparing said level to a standard,
 wherein a reduced level of said peptide is indicative of said magnesium binding defect.
- 12 (Original) The method of claim 11, wherein said level of peptide is measured by using an antibody to peptide having an amino acid sequence selected from the group consisting of: Phe-Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:1), Phe-Xaa-Gly-Leu-Met-NH₂, where Xaa is variant Phe or Val (SEQ ID NO:4), and Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:2).
 - 13 (Original) The method of claim 12, wherein the antibody is monoclonal.
- (Original) The method of claim 13, wherein the monoclonal antibody cross reacts with each of said peptides
- 15 (Original) The method of claim 12, wherein the antibody is employed in an immunoenzyme assay.

- (Original) The method of claim 15, wherein the immunoenzyme assay is enzyme-linked immunosorbent assay to quantitate the concentration of said peptide in blood serum.
 - 17 (Original) The method of claim 12, wherein the antibody is polyclonal.
 - 18 (Cancelled)
 - (Cancelled)
 - 20 (Cancelled)
- 21 (New) A method for monitoring progress in treatment of the magnesium binding defect in an individual, comprising:
 - a) measuring the level of peptide in a sample of body fluid of said individual, said peptide having an amino acid sequence selected from the group consisting of: Phe-Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:1), Phe-Xaa-Gly-Leu-Met-NH₂, where Xaa is variant Phe or Val (SEQ ID NO:4), and Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:2); and
- b) comparing said level of peptide to the level of the peptide after treatment, whereby a significant increase in the level of said peptide is indicative of progress of treatment of said individual.
- 22 (New) The method of claim 21, wherein said level of peptide is measured by using an antibody to peptide having an amino acid sequence selected from the group consisting of: Phe-Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:1), Phe-Xaa-Gly-Leu-Met-NH₂, where Xaa is variant Phe or Val (SEQ ID NO:4), and Phe-Gly-Leu-Met-NH₂ (SEQ ID NO:2).
 - 23 (New) The method of claim 22, wherein the antibody is monoclonal.